SEQUENCE LISTING

<110> Edgar B. Cahoon Rebecca E. Cahoon William D. Hitz Anthony J. Kinney <120> Membrane-Bound Desaturases <130> BB1264 ` <140> US/09/857,524 <141> 2001-06-04 <150> 60/110,784 <151> 1998-12-03 <160> 17 <170> Microsoft Office 97 <210> 1 <211> 1471 <212> DNA <213> Picramnia pentandra <220> <221> unsure <222> (1402) <223> n = A, C, G, or T <400> 1 cttccttgtt cctggaattt tcaaatcact tcctctgttg cacttcaatg gaagagccaa 60 agaagcacat ttcgcaagca gaccttgcaa agcataagca accaggagat ttatggatct 120 ctatcaaggg aaaagtttac gatatctcca agtggactaa agagcatccc ggtggtgagc 180 tcccattgtt aagttttgcc ggccaagatg tcactgatgc gttcattgct taccatcctg 240 300 gcactgcttg gcaatacctt gacaggttct ttactgggta ctacgttcaa gattactctg tetetgagat gtecaaggae tacagaagge tegtetetga gttttetaag atgggtttgt 360 tcaagacacc aggcaaaggg gtctactgct caatcttttt cgtgtctgtg ttgttcgctc 420 tgagtgttta cggtgttctc tactgcaaga gcacctgggc tcatctttgc tctggtttgc 480 taatgggtat gctatggctc cagagtggtt gggtggggca tgattcttgt cactaccaag 540 ttatgcctaa ccgtaagctt aatcgtcttt ttcaaatcat tgcaggaaat gtgattgctg 600 gtgttagtgt tgcatggtgg aagttggacc ataacaccca tcactttgcc tgtaatagcg 660 720 ccaatctqga tcctgatatt cagcacette ctataattge catateecca aaatttttea 780 actecettae ateataetat cacaaetgea aaatgaeeta tgategeget geeaggtttt ttgttagctt tcagcactgg acattttatc ctgcattgtt aagcgttagg ctctatcttt 840 ttattctgtc ttttaaggtg gtgttttcca acaacaaaag ggtatacaag agaagtcagg 900 aaattttagg ctatgcagct ttcttgactt ggtattctct actcctttct cgcctaccca 960 attggcctga aagggtcatg tatttcacgt cctgtttagc agtcgccggg ttccaacatt 1020 ggcaqttcag cttgaatcac tttgcttcta atgtttacac tggtttgcct agcggtaatg 1080 attggtttca ccagcagaca aagggcacgc tcaacataac agcttctgct tggtgggatt 1140 ggtttcatgg tggcctgcac tttcagattg agcatcatct gtttccaagg atgcctaagt 1200 gccatttcag gaaaatctca cccattgtga acaaactttg ccagaagcat aatttgtcct 1260 atgaaactgc taccatgtgg gaggccaata aaatggtata ctccaccctg cgtgctgtgg 1320 ctatqqaaqc taaqqatqtt accaaqccag ttcccaagaa catggtctgg gaagcaatga 1380 acactttcgg gtgaacctta tnaaacatca agtgctgtct ttcccgtaaa agcttccagt 1440 cccaatgttt ctttttttt tttttttt t 1471 <210> 2

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<213> Picramnia pentandra

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305					310					315					320	
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Glu Lys Arg Val Pro Gln Arg Leu Leu Glu Ile Ala Gly Val Ala Thr 295 300 290 Phe Trp Ala Trp Tyr Pro Leu Leu Val Ala Ser Leu Pro Asn Trp Trp 305 310 315 320 Glu Arg Val Ala Phe Val Leu Phe Ser Phe Thr Ile Cys Gly Ile Gln 325 330 335 His Val Gln Phe Cys Leu Asn His Phe Ser Ser Asp Val Tyr Val Gly 340 345 350 Pro Pro Lys Gly Asn Asp Trp Phe Glu Lys Gln Thr Ala Gly Thr Leu 360 355 365 Asp Ile Leu Cys Ser Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln 370 375 380 Phe Gln Ile Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu 395 400 385 390 Arg Lys Val Ala Pro Ala Val Arg Asp Leu Cys Lys His Gly Leu 405 410 415 Thr Tyr Ser Ala Ala Thr Phe Trp Gly Ala Asn Val Leu Thr Trp Lys 425 430 420 Thr Leu Arg Ala Ala Ala Leu Gln Ala Arg Thr Ala Thr Ser Gly Gly 435 440 445 Ala Pro Lys Asn Leu Val Trp Glu Ala Val Asn Thr His Gly 455 460 450 <210> 5 <211> 880 <212> DNA <213> Glycine max · <220> <221> unsure <222> (496) <223> n = A, C, G, or T <220> <221> unsure <222> (512) <223> n = A, C, G, or T <220> <221> unsure <222> (523) <223> n = A, C, G, or T <220> <221> unsure <222> (532) <223> n = A, C, G, or T <220> <221> unsure

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Ser Thr Phe Val His Val Leu Ser Ala Ala Leu Ile Gly Phe Leu Trp Ile Gln Ser Gly Trp Ile Gly His Asp Ser Gly His Tyr Asn Val Met Leu Ser Arg Arg Leu Asn Arg Ala Ile Gln Ile Leu Ser Gly Asn Ile Leu Ala Gly Ile Ser Ile Gly Trp Trp Lys Trp Asn His Asn Ala His His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His Met Pro Val Phe Ala Val Ser Ser Arg Phe Phe Asn Ser Ile Thr Ser His Xaa Tyr Gly Arg Lys Xaa Glu Phe Asp Xaa Ile Ala Xaa Phe Leu Ile Cys Tyr Gln His Phe Thr Phe Tyr Pro Val Met Cys Val Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Phe Ser Arg Xaa Lys Val Gln Asp Arg Ala Leu Asn Ile Met Gly Ile Leu Val Phe Trp Thr Trp Phe Leu Phe Leu Leu Ala Leu Leu Phe Val Pro Ile Gln His Ile Gln Phe Trp Leu Asn His Leu Ala Glu Asn Leu Tyr Xaa Gly <210> 7 <211> 1934 <212> DNA <213> Glycine max <400> 7 gcacgagcac acaagtaaaa ccttagagag agagagagag agagagagag ggtaaaaggg tattagatcc ttgaaccaga tcaaatcatc aaaatctctg tctatggggt tgtgaaaaca 120 gaagaaccag ataccccatt attgttctta tctatctatc tgtctatatt ctatttatct tottattgta gttotcattg tgtotgattt cagtgatttg tgttgttttt ggttaacaca agcaatgqag gttgttgaga aggagaagaa gtacataacc tcagaggagc tgaagggtca caacaaqqaq qqaqatttat ggatctcaat tcaaggtaag gtgtacaatg tctcagattg ggtcaaggag caccetggtg gtgatgttee aateteaaae ettgetggee aggatgteae tgatgcattc atagcatacc atcctggcac agcatggtca caccttgaaa aattcttcac 540 tggctaccac ctcagtgact tcaaggtctc tgaggtgtcc aaagactaca gaaagcttgc 600 atctgagttc tcaaaattgg gtctttttga caccaaaggg catgtcactt catgcaccct tgcatctgtt gctgttatgt tcctcattgt actctatggt gttctgaggt gcactagtgt gtgggctcat ttgggttcag gcatgctctt agggttgctt tggatgcaaa gtgcttatgt gggccatgat tctggccact atgtggttat gacaaccaat ggtttcaaca aggttgcaca gatcctctct gggaactgct tgaccgggat aagcattgct tggtggaagt ggactcacaa tgctcaccac attgcgtgca acageettga ceatgaeeet gatetgeage acatgeeggt ctttgcagtt tcgtcgcggt tcttcaattc cataacctct catttctatg ggaggaagtt 1020 ggagtttgat ttcattgcta ggttcttgat ctgctaccag cactttactt tttacccggt 1080





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Asp His Asp Pro Asp Leu Gln His Met Pro Val Phe Ala Val Ser Ser Arg Phe Phe Asn Ser Ile Thr Ser His Phe Tyr Gly Arg Lys Leu Glu . Phe Asp Phe Ile Ala Arg Phe Leu Ile Cys Tyr Gln His Phe Thr Phe Tyr Pro Val Met Cys Val Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Phe Ser Arg Arg Lys Val Gln Asp Arg Ala Leu Asn Ile Met Gly Ile Leu Val Phe Trp Thr Trp Phe Pro Leu Leu Val Ser Cys Leu Pro Asn Trp Pro Glu Arg Val Met Phe Val Leu Ala Ser Phe Ala Val Cys Ser Ile Gln His Ile Gln Phe Cys Leu Asn His Phe Ala Ala Asn Val Tyr Val Gly Pro Pro Ser Gly Asn Asp Trp Phe Glu Lys Gln Thr Ser Gly Thr Leu Asp Ile Ser Cys Ala Ser Ser Met Asp Trp Phe Phe Gly Gly Leu Gln Phe Gln Leu Glu His His Leu Phe Pro Arg Leu Pro Arg Cys Gln Leu Arg Lys Ile Ser Pro Leu Val Ser Asp Leu Cys Lys Lys His Asn Leu Pro Tyr Arg Ser Leu Ser Phe Trp Glu Ala Asn Gln Trp Thr Ile Arg Thr Leu Arg Thr Ala Ala Leu Gln Ala Arg Asp Leu Thr Asn Pro Ala Pro Lys Asn Leu Leu Trp Glu Ala Val Asn Thr His Gly <210> 9 <211> 1972 <212> DNA <213> Triticum aestivum <400> 9 gcacgagete cetaacaaac etcegttget gttttaagat eegateteee etteeeeet cccctccctt cctcctgagt cctgaccacc cctcctcgcg ctccagctaa atccacgcca ccgatggccc gcacgggcct cgcggacgca acggcgccgg aagccgacgc aatgccggcc gccagcaagg acgccgccga cgtccgcatg atctccacca aggagctgca ggcgcacgct gccgcggacg acctctggat ctccatctcc ggggacgtct acgacgtcac gccgtggctg cgccaccacc cgggcggcga ggtcccgctc atcaccctcg ccggccagga cgccaccgac 360 gccttcatgg cctaccaccc gccctccgtg cgcccgctcc tccgccgctt cttcgtcggc 420

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Met Phe Ala Gly Gly Leu Ile Gly Phe Ile Trp Ile Gln Ser Gly Trp Ile Gly His Asp Ser Gly His His Gln Ile Thr Arg His Pro Ala Leu Asn Arg Leu Leu Gln Val Val Ser Gly Asn Cys Leu Thr Gly Leu Gly Ile Ala Trp Trp Lys Phe Asn His Asn Thr His His Ile Ser Cys Asn Ser Leu Asp His Asp Pro Asp Leu Gln His Leu Pro Leu Phe Ala Val Ser Thr Lys Leu Phe Asn Asn Leu Trp Ser Val Cys Tyr Glu Arg Thr Leu Ala Phe Asp Ala Ile Ser Lys Phe Phe Val Ser Tyr Gln His Trp Thr Phe Tyr Pro Val Met Gly Phe Ala Arg Ile Asn Leu Leu Val Gln Ser Ile Val Phe Leu Ile Thr Gln Lys Lys Val Arg Gln Arg Trp Leu Glu Ile Ala Gly Val Ala Ala Phe Trp Val Trp Tyr Pro Leu Leu Val Ser Cys Leu Pro Asn Trp Trp Glu Arg Val Ala Phe Val Leu Ala Ser Phe Val Ile Thr Gly Ile Gln His Val Gln Phe Cys Leu Asn His Phe Ser Ser Ala Val Tyr Val Gly Pro Pro Lys Gly Asn Asp Trp Phe Glu Arg Gln Thr Ala Gly Thr Leu Asp Ile Lys Cys Ser Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln Phe Gln Val Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Tyr Arg Met Val Ala Pro Ile Val Arg Asp Leu Cys Lys Lys His Gly Leu Ser Tyr Gly Ala Ala Thr Phe Trp Glu Ala Asn Val Met Thr Trp Lys Thr Leu Arg Ala Ala Ala Leu Gln Ala Arg Glu Ala Thr Thr Gly Ala Ala Pro Lys Asn Leu Val Trp Glu Ala Leu Asn Thr His Gly



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Cys Leu Val Phe Ser Ile Trp Tyr Pro Leu Leu Val Ser Cys Leu Pro Asn Trp Gly Glu Arg Ile Met Phe Val Ile Ala Ser Leu Ser Val Thr Gly Met Gln Gln Val Gln Phe Ser Leu Asn His Phe Ser Ser Val Tyr Val Gly Lys Pro Lys Gly Asn Asn Trp Phe Glu Lys Gln Thr Asp Gly Thr Leu Asp Ile Ser Cys Pro Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln Phe Gln Ile Glu His His Leu Phe Pro Lys Met Pro Arg Cys Asn Leu Arg Lys Ile Ser Pro Tyr Val Ile Glu Leu Cys Lys His Asn Leu Pro Tyr Asn Tyr Ala Ser Phe Ser Lys Ala Asn Glu Met Thr Leu Arg Thr Leu Arg Asn Thr Ala Leu Gln Ala Arg Asp Ile Thr Lys Pro Leu Pro Lys Asn Leu Val Trp Glu Ala Leu His Thr His Gly <210> 12 <211> 469 <212> PRT <213> Triticum aestivum <400> 12 Met Ala Arg Thr Gly Leu Ala Asp Ala Thr Ala Pro Glu Ala Asp Ala Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr Lys Glu Leu Gln Ala His Ala Ala Ala Asp Asp Leu Trp Ile Ser Ile Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe



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Cys Ile Ala Leu Tyr Cys Val Leu Ala Cys Ser Ser Thr Gly Ala His Met Phe Ala Gly Gly Leu Ile Gly Phe Ile Trp Ile Gln Ser Gly Trp Ile Gly His Asp Ser Gly His His Gln Ile Thr Arg His Pro Ala Leu Asn Arg Leu Leu Gln Val Val Ser Gly Asn Cys Leu Thr Gly Leu Gly Ile Ala Trp Trp Lys Phe Asn His Asn Thr His His Ile Ser Cys Asn Ser Leu Asp His Asp Pro Asp Leu Gln His Leu Pro Leu Phe Ala Val Ser Thr Lys Leu Phe Asn Asn Leu Trp Ser Val Cys Tyr Glu Arg Thr Leu Ala Phe Asp Ala Ile Ser Lys Phe Phe Val Ser Tyr Gln His Trp Thr Phe Tyr Pro Val Met Gly Phe Ala Arg Ile Asn Leu Leu Val Gln Ser Ile Val Phe Leu Ile Thr Gln Lys Lys Val Arg Gln Arg Trp Leu Glu Ile Ala Gly Val Ala Ala Phe Trp Val Trp Tyr Pro Leu Leu Val Ser Cys Leu Pro Asn Trp Trp Glu Arg Val Ala Phe Val Leu Ala Ser Phe Val Ile Thr Gly Ile Gln His Val Gln Phe Cys Leu Asn His Phe Ser Ser Ala Val Tyr Val Gly Pro Pro Lys Gly Asn Asp Trp Phe Glu Arg Gln Thr Ala Gly Thr Leu Asp Ile Lys Cys Ser Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln Phe Gln Val Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Tyr Arg Met Val Ala Pro Ile Val Arg Asp Leu Cys Lys His Gly Leu Ser Tyr Gly Ala Ala Thr Phe Trp Glu Ala Asn Val Met Thr Trp Lys Thr Leu Arg Ala Ala Ala Leu Gln Ala Arg Glu Ala Thr Thr Gly Ala Ala Pro Lys Asn Leu Val Trp Glu Ala



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۹. Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Ile Ser Lys Arg Lys 280 285 275 Ile Pro Asp Arg Gly Leu Asn Ile Leu Gly Thr Leu Ile Phe Trp Thr 290 295 300 Trp Phe Pro Leu Leu Val Ser Arg Leu Pro Asn Trp Pro Glu Arg Val 305 310 315 320 Ala Phe Val Leu Val Ser Phe Cys Val Thr Gly Ile Gln His Ile Gln 325 330 335 Phe Thr Leu Asn His Phe Ser Gly Asp Val Tyr Val Gly Pro Pro Lys 340 345 350 . Gly Asp Asn Trp Phe Glu Lys Gln Thr Arg Gly Thr Ile Asp Ile Ala 355 360 365 Cys Ser Ser Trp Met Asp Trp Phe Phe Gly Gly Leu Gln Phe Gln Leu 375 380 370 Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu Arg Ser Ile 395 400 390 385 Ser Pro Ile Cys Arg Glu Leu Cys Lys Tyr Asn Leu Pro Tyr Val 405 410 415 Ser Leu Ser Phe Tyr Asp Ala Asn Val Thr Thr Leu Lys Thr Leu Arg 425 430 420 Thr Ala Ala Leu Gln Ala Arg Asp Leu Thr Asn Pro Ala Pro Gln Asn 445 435 440 Leu Ala Trp Glu Ala Phe Asn Thr His Gly 450 455 <210> 14 <211> 35 <212> DNA <213> Artificial Sequence <220> <223> Definition of Artificial Sequence: PCR primer for 5' of pk0011.d5 <400> 14 tttgcggccg caaatcaatg gaagaagcaa agaag 35 <210> 15 <211> 33 <212> DNA <213> Artificial Sequence <220> <223> Definition of Sequence: antisense PCR primer for 3' of pk0011.d5 <400> 15 tttgcggccg ccaggattca cccgaaagtg ttc <210> 16 <211> 823 <212> DNA



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